VERSION WITH MARKINGS TO SHOW CHANGES MADE

In the Claims:

- 2. (Amended) The array according to Claim 1, wherein two or more different target nucleic acids are represented hybridize to different probe oligonucleotide spots in said pattern.
- 8. (Amended) The array according to Claim 7, wherein said-each of said long oligonucleotide probes is cross-linked to the surface of said support at at least one site.
- 15. (Amended) The array according to Claim 14, wherein <u>said array comprises</u> ten or more different <u>target nucleie acids are represented probe oligonucleotide spots</u> in said pattern, each of which <u>hybridizes to a different target nucleic acid</u>.
- 36. (Amended) The array according to Claim 1, wherein any variance in hybridization efficiency among any two probes of said array does not exceed about 10-fold as determined using the hybridization efficiency assay described in the specification beginning at page 23, lines 4 ff.
- 37. (Amended) The array according to Claim 14, wherein any variance in hybridization efficiency among any two probes of said array does not exceed about 10-fold as determined using the hybridization efficiency assay described in the specification beginning at page 23, lines 4 ff.
- 38. (Amended) The array according to Claim 23, wherein any variance in hybridization efficiency among any two probes of said array does not exceed about 10-fold as determined using the hybridization efficiency assay described in the specification beginning at page 23, lines 4 ff.

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